



Article

Characterization and Bioactivity of Polysaccharides Separated through a (Sequential) Biorefinery Process from *Fucus spiralis* Brown Macroalgae

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Abstract: Marine macroalgae biomass is a valuable renewable resource that can be used for the development of bioeconomy through the valorisation of valuable compounds. The aim of the current study is separate macroalgal polysaccharides with bioactive properties from brown macroalgae *Fucus spiralis* based on a designed biocascading biorefinery approach. Thus, we applied an integrated processing method for the separation of fucoidan and alginate, in addition to characterization through IR spectroscopy and ¹H NMR. The bioactivity potential (antioxidant activity using superoxide anion and DPPH radical scavenging analysis) of the two polysaccharides was evaluated, together with DNA binding studies performed through voltametric techniques and electronic spectroscopy titration. In terms of results, functional groups S=O (1226 cm⁻¹), N=S=O (1136 cm⁻¹) and C-O-SO₃ (1024 cm⁻¹), which are characteristic of fucoidan, were identified in the first polysaccharidic extract, whereas guluronic units (G) (1017 cm⁻¹) and mannuronic units (M) (872 and 812 cm⁻¹) confirmed the separation of alginate. The DNA binding studies of the isolated polysaccharides revealed an electrostatic and an intercalation interaction of DNA with fucoidan and alginate, respectively. Both antioxidant activity assays revealed improved antioxidant activity for both fucoidan and alginate compared to the standard α-tocopherol.

Keywords: brown algae; fucoidan; alginate; antioxidants; DNA binding study; bioactivity



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1. Introduction

Marine macroalgae, also known as “seaweeds”, represent a promising biomass that can address future trends and industry demand, as well as various sustainability issues. Marine macroalgae have been recognized as an alternative for the production of renewable energy, bio-based products and bioactive molecules, owing to their remarkable regeneration properties and high photosynthetic efficiency, as well as the lack of land requirements for growth [1–4]. Yields in terms of growth potential of seaweeds are around 20 t per hectare per year [5]. Furthermore, the valorisation of invasive marine macroalgae could provide additional advantages through processing [6], whereas others are suitable for cocultivation with fish species in an integrated multitrophic aquaculture system (IMTA), where they play a bioremediation role [7]. The chemical composition of the marine macroalgae, including the bioactivity of valuable compounds, depends on the harvesting season, the algae species and the environmental conditions [8]. Furthermore, the separation method has a significant influence on the final chemical structure of the targeted compounds [9].

Processing of marine macroalgae through a biorefinery approach could enable the generation of cost-effective products and a sustainable valorisation with minimal waste or even the fulfilment of the zero-waste concept [10–13]. The amount of easily degradable carbohydrates in macroalgae typically ranges between 25 and 60% dry wt [5]. The lack

of or minimal amount of lignin in macroalgae facilitates separation of carbohydrates compared with terrestrial biomass [14]. Therefore, their chemical composition makes marine macroalgae suitable for generation of fermentation products, such as bioethanol, methane, biohydrogen, bio-oil and biodiesel [5], as well as valuable bioactive molecules.

Brown macroalgae have been traditionally used as food and medicine in Asian countries. Globally, they are mainly applied as feed and crop fertilizer, owing to their high ash and mineral content [14,15]. Additionally, this biomass is highly complex in its chemical structure and includes valuable bioactive metabolites, some of which are unique in the natural world. Polysaccharides represent the main constituent of the cell wall of marine macroalgae and have important bioactive properties. Fucoidan, as well as some polyphenols, from brown algae can be used for the production of pharmaceuticals, nutraceuticals and cosmetics, owing to their high bioactivity potential [16–18]. Alginate, a macroalgal hydrocolloid, is mainly used in food, pharmaceutical and medical applications, as well as in textile and paper production [19–21]. Furthermore, its gelling and thickening properties have been increasingly exploited to generate innovative biofilms and hydrogels [22,23]. The separation of bioactive compounds is recommended to be performed in the initial steps of the biorefining process in order to ensure the feasibility of the process [24,25].

Fucus spiralis, which belongs to the order *Fucales*, is an edible and perennial brown macroalga common throughout the temperate regions of the northern hemisphere [26,27] occupying the littoral to sublittoral areas of rocky coasts [28]. Its chemical composition includes proteins with antioxidant properties and angiotensin-converting enzyme (ACE) inhibitory activity [29]; polyphenols, such as phlorotannins with antioxidant activity [30–32]; and anti-inflammatory properties [33]; as well as polysaccharides with anticoagulant [34], anticancer [35,36] and antioxidant activity [37].

Sequential processes applied to date for biorefining of brown algae biomass have consisted of the coproduction of bioactive compounds and/or biofuel, mostly from species of the genera *Laminaria* [15,21,38–41], *Sargassum* [13,42,43], *Macrocystis* [44,45], *Ascophyllum* [15] and *Ecklonia* [44,46].

The purpose of this work is (i) to design and perform an integrated valorisation of *Fucus spiralis* biomass, applying soft or green processes for the separation of high-value molecules; and for the targeted compounds, (ii) the evaluation of biological potential through DNA binding studies (an electrochemical DNA interaction study and an electronic spectroscopy DNA interaction assay) and antioxidant activity studies (an superoxide anion radical interaction study and a DPPH assay). To the best of our knowledge, no studies have been conducted on a biorefinery approach for *F. spiralis* macroalgae, and no evidence of the bioactive properties of the targeted molecules have been reported to date.

Therefore, in the current study, we highlight a pathway that can be applied in order to process brown macroalgae *Fucus spiralis* through a biorefinery approach, in addition to reporting the bioactive properties of the obtained biocompounds.

2. Materials and Methods

Fucus spiralis algae were collected during the winter season in the Viana do Castelo area, on the northern coast of Portugal. The biomass was washed with tap water, followed by a few washes with distilled water in order to eliminate impurities, sand and epiphytes. Algal biomass was further prepared by drying in an oven at 50 °C for 24 h. The dried algae were ground into particles with a size ≤ 0.5 cm and kept in a desiccator in sealed bags.

All reagents and solvents were of analytical grade and were used without further purification. Dimethyl sulfoxide (DMSO) and ethanol (HPLC-grade; Sigma-Aldrich, St. Louis, MO, USA) were used as solvents in voltametric and spectroscopic assays, whereas tetrabutylammonium tetra-fluoroborate (Bu_4NBF_4) (electrochemical grade 99%; Sigma-Aldrich) was used as a supporting electrolyte. Research-grade nitrogen and oxygen gases (99.99%) were provided by Linde Gas Algérie.

2.1. Methods of Extraction

2.1.1. Extracts Containing Polyphenols

Polyphenols were extracted with 70% ethanol as solvent in a batch sonication system (Elmasonic 120 (H), Lauda Konigshofen, Germany; power, 80–320 W; frequency, 35 kHz), using a solid to liquid ratio of 1:10 at 40 °C, with an extraction time of 45 min [47]. The crude extract was filtered, and the residual biomass was dried at 40 °C and directly used for further extraction steps. The final filtered extract containing polyphenols is hereafter abbreviated as E₁.

2.1.2. Extracts Containing Polysaccharides

The two main macroalgal polysaccharides, fucoidan and alginate, were sequentially extracted from the obtained residue after the separation of polyphenols by adapting a protocol reported by Yuan and Macquarrie [48]. First, fucoidan was extracted using HCl 0.1 M as solvent, with a solid/liquid ratio of 1:10, at 40 °C, with a 20 min extraction time.

The alginate was isolated using a cost-effective and soft separation method with 4% Na₂CO₃ solution. Both extracts were filtered and precipitated with a 1:1 *v/v* ethanol:extract ratio. The residual biomass and the filtered extracts were washed with distilled water and dried at 40 °C. The final extracts are hereafter abbreviated as E₂ (fucoidan) and E₃ (alginate).

2.2. Chemical Characterization of Extracts

In order to determine the total polyphenolic content (TPC), E₁ analysed according to the Folin–Ciocalteu method and quantified in mg gallic acid equivalents per gram of biomass (mg GAE g⁻¹) [49].

We evaluated the total reducing sugar content in the supernatant produced after the separation and filtration of the E₂ and E₃ extracts, as well as in the wastewater resulting from the purification processes. The concentration of the reducing sugars was determined following the method described by the IUPAC [50], using an Avanta UV-VIS GBS spectrophotometer at 540 nm.

Elemental analysis was carried out using the Pregl method [51] for carbon and hydrogen, the Kjeldahl method [52] for nitrogen and the Schöniger method [53,54] for sulphur, with all results expressed in percentage.

The FTIR spectra were obtained in duplicate using an IRAffinity-1 Shimadzu FTIR instrument in the wavenumber range of 400–4000 cm⁻¹.

Proton nuclear magnetic resonance (¹H NMR) spectra were recorded by a Bruker Avance DRX 400 MHz spectrometer using deuterated water as solvent and tetramethylsilane as internal reference. Chemical shifts are reported in parts per million (ppm) relative to the residual peak of the deuterated water used as solvent. For the preparation of samples, 5 mg each of dry extracts E₂ and E₃ was dissolved in 800 µL of D₂O.

2.3. Determination of Extract Bioactivity

2.3.1. DNA Binding Studies

Double-stranded DNA (ds. DNA) was extracted from chicken blood using a nuclear cell lysis method [55]. Absorption spectroscopy and the molar absorption coefficient value of 6600 M⁻¹·cm⁻¹ at 260 nm were used to determine the concentration of DNA [56]. The obtained ratio of absorbance of A₂₆₀/A₂₈₀ in the DNA sample was 1.97, reflecting the purity of extracted ds. DNA, which was apparently free from proteins. All stock solutions were used within 5 days after preparation and stored at 4 °C until use.

Voltametric assays were performed using a PGZ301 voltammeter running on VoltMaster 4 V 7.08 software, (Radiometer Analytical SAS, Lyon, France). The concentration of the supporting electrolyte was kept at 0.1 M. Nitrogen gas was bubbled through the solution to avoid any interference from air. Experiments were carried out in a 12 mL three-electrode electrochemical cell containing a platinum disk (Pt) working electrode with a geometric area of 0.013 cm², a platinum wire as counter (auxiliary) electrode and an Hg/Hg₂Cl₂ paste-covered wire as reference electrode.

As a complementary method to the voltammetry techniques, electronic spectroscopy titration was performed to study the binding affinity of E₂ and E₃ with DNA in buffer phosphate solution (KH₂PO₄/K₂HPO₄) at pH = 7.2. The absorption spectra of a fixed amount of 2 mg of E₂ and 6 mg of E₃ were recorded in the absence and presence of a gradually increasing concentration of DNA stock solution. Absorption spectra measurements were conducted on a UV-Vis spectrometer, (Shimadzu 1800, Japan).

2.3.2. Superoxide Anion Radical Assay

Cyclic voltammetry measurements were performed on a platinum electrode at 100 mV·s⁻¹ using a PGZ301 voltammeter running on VoltaMaster 4 V 7.08 software (Radiometer Analytical SAS, France).

In order to obtain kinetic curves and to calculate the IC₅₀ values, the O₂^{·-} radical-scavenging activity was plotted based on Equation (1) against varying compound concentrations [57–59]. The antioxidant capacity of E₂ and E₃ was expressed as IC₅₀. The IC₅₀ value was defined as the concentration (mg/mL) of samples that inhibited the formation of O₂^{·-} radicals by 50%.

$$\% \text{ radical scavenging activity} = \frac{i_{pa0} - i_{pa}}{i_{pa0}} \times 100 \quad (1)$$

where i_{pa0} and i_{pa} are the anodic peak current densities of the superoxide anion radical in the absence and presence of E₂ and E₃, respectively.

2.3.3. DPPH (2,2-Diphenyl-1-picrylhydrazyl) Assay

Spectrophotometric experiments were performed to measure the antioxidant activity of E₂ and E₃ samples using DPPH free radical. Experiments were carried out by adding gradually increasing concentrations of E₂ in DMF (5 mg/mL) or E₃ (5 mg/mL) to a 2.1 mM DMF solution of DPPH. Absorbance at 524 nm was measured after 30 min of incubation in the dark. Finally, Equation (1) was applied to calculate % inhibition of DPPH. In order to obtain kinetic curves and to calculate the IC₅₀ values, DPPH radical-scavenging was plotted against varying compound concentrations. The antioxidant capacity of E₂ and E₃ was expressed as IC₅₀.

3. Results

3.1. Biomass Proximate and Ultimate Analyses

Prior to designing a biorefinery approach with the goal of recovering valuable molecules, the feedstock (*Fucus spiralis* macroalgae) was subjected to proximate and ultimate analyses; the chemical composition is listed in Table 1. The *Fucus spiralis* ash content is similar to the values reported in the literature for other algae species: an ash content of 21.0 ± 0.2% dw and 28.0 ± 0.2% dw for *Laminaria digitata* and 25.0 ± 0.2% dw and 26.5 ± 0.7% dw for *Fucus spiralis* [60]. Furthermore, [61] we determined an ash percentage of 31.0 ± 0.1% dw for untreated *Laminaria digitata*, whereas the washed alga generated a value of 7.9 ± 0.1% dw.

The amount of reducing sugars detected in *Fucus spiralis* biomass was 16.3%. Lower limits of 9.71 ± 0.03% dw [62] and 10–11% dw [28] were reported, as well as higher concentrations [63]. With respect to the amount of proteins amount, the percentage obtained in the current study for *Fucus spiralis* is close to values registered in the case of *Fucus vesiculosus*, i.e., 12.99% dw [64] or 11% dw [28].

Table 1. Chemical characteristics of main *Fucus spiralis* biomass.

Property Value	Unit	Value
Proximate analysis		
Moisture content	wt% (dry)	11.5 ± 0.7
Ash content	wt% (dry)	22.5 ± 0.9
Ultimate analysis		
Carbon	wt% (dry)	36.6 ± 0.6
Hydrogen	wt% (dry)	3.9 ± 0.1
Nitrogen	wt% (dry)	1.1 ± 0.1
Sulphur	wt% (dry)	0.5 ± 0.1
Carbohydrate analysis		
Reducing sugars	wt% (dry)	16.3 ± 0.7
Protein characterization		
Protein content	wt% (dry)	5.7 ± 0.5
Nitrogen-to-protein conversion factor	-	6.25

3.2. A Biorefinery Approach for *Fucus spiralis* Macroalgal Biomass Processing

Based on the chemical characterization of the feedstock and available studies concerning sequential separation of macroalgal compounds [39,44,48,65], a biorefinery flow sheet for *Fucus spiralis* biomass was designed (Figure 1). All solvents and methods were selected taking into consideration cost effectiveness and circular economy. Thus, a sustainable, efficient and minimal-waste approach was considered. The primary biorefinery step was designed to obtain easily extractible valuable molecules, such as polyphenols and saccharides.

The sequence starts with green light sonication (Figure 1) in order to separate the polyphenols. This step could also act as a pretreatment to make the biomass more suitable for the following extraction steps. Polyphenol separation in the initial biorefinery step is recommended in order to avoid the contamination of further extracted molecules [46,66,67]. Furthermore, *Fucus* is among the brown macroalgae genera with the highest percentage of polyphenolic compounds (1–14% dw) [31]. Usually, polyphenol separation from marine macroalgae is performed using formaldehyde as, as polyphenols have tight chemical bonds with macroalgal polysaccharides [68,69]. However, formaldehyde is known to be a carcinogenic solvent. Therefore, polyphenol extraction was performed with ethanol (70%), considering the positive yields it provides, GRAS recognition (Generally recognized as safe) and low cost [70,71]. Polyphenols were extracted in a batch sonication system using a 1/10 solid/liquid ratio at 40 °C, with an extraction time of 45 min.

For the extraction of fucoïdan, hydrochloric acid was selected as solvent, as it has shown satisfactory results in extracting fucoïdan in single processing or even in sequential extractions from brown algae [48]. After washing the algal residue in distilled water, alginate is the next compound to be extracted using sodium carbonate [19,44]. A low temperature was chosen in order to avoid the degradation of the targeted metabolites [28,67]. Fucoïdan and alginate must be separated sequentially; otherwise, both molecules would be separated in the same extract following the degradation of the algal cell wall [72].

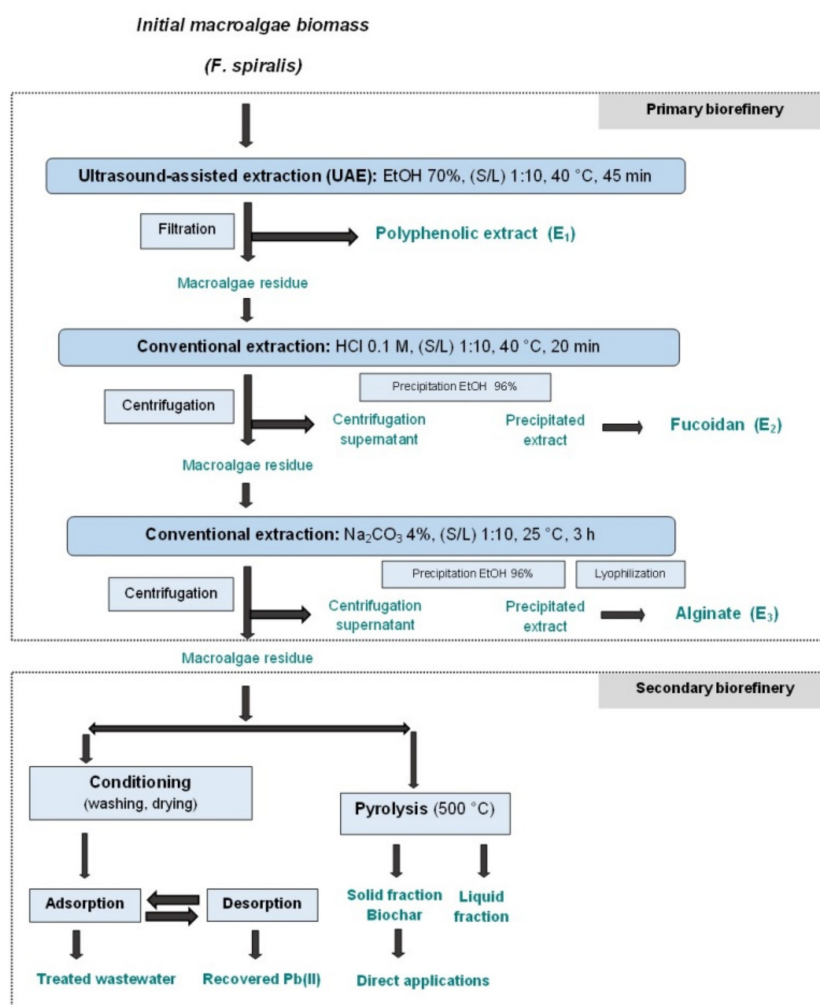


Figure 1. The biorefinery flow sheet designed for *Fucus spiralis* biomass.

The *Fucus spiralis* solid residue generated as waste after the separation of bioactive molecules was designated as a biosorbent for a cascade of sorption/desorption cycles. Several studies have dealt with the use of macroalgae to uptake pollutants from wastewater due to their effectiveness in the removal of low concentrations and relatively low cost in comparison to conventional adsorbents [73–77]. The literature has proven that brown macroalgae are very good biosorbents and are more effective in the removal of harmful pollutants than red and green algae [78,79]. In most previous studies, including research using *Fucus spiralis* [73], researchers used the entire algal biomass as biosorbent, whereas few studies involved biosorption tests with seaweed waste. Examples include the use of red algae waste generated from agar extraction to adsorb Cu^{+2} , Pb^{+2} and Cd^{+2} [80–82] and the application of dealginated brown macroalgae waste [83,84]. Residues derived from brown macroalgae *Laminaria digitata* and green macroalgae *Ulva lactuca* were used as biosorbent for Ce^{+3} , Pb^{+2} , Cu^{+2} and Ni^{+2} [85]. However, only our team has reported satisfactory biosorption results with *Fucus spiralis* waste [75].

Considering all these aspects, the designed biorefinery approach was regarded to be a viable option for the processing of *Fucus spiralis* biomass and was therefore applied in the current study.

3.3. Extract Characterization

3.3.1. Yield and Chemical Characterization of Polyphenolic Extract (E₁)

The *Fucus spiralis* polyphenol extract E₁ registered a total polyphenolic content (TPC) of 3.4 ± 0.0 mg GAE g⁻¹. This value is lower than those obtained in [86] for *Laminaria digitata* (37.66 ± 0.00 mg GAE g⁻¹ dry seaweed) and *Laminaria saccharina* (66.75 ± 3.72 mg GAE g⁻¹ dry seaweed) collected from Northern Ireland but comparable to those generated for conventional polyphenol separation from *Laminaria japonica* (38.47 ± 0.88 GAE mg/100 g dry seaweed), *Lessonia trabeculate* (49.80 ± 5.68 GAE mg/100 g dry seaweed) and *Ascophyllum nodosum* (51.47 ± 3.28 GAE mg/100 g dry seaweed) [87]. The geographic areas in which the algae originated, as well as the time of collection and pretreatment processes, could explain these differences. The polyphenol extraction yield obtained from *Fucus spiralis* was 14.0% dry seaweed.

3.3.2. Yield and Chemical Characterization of Fucoïdan Extract (E₂)

Fucoïdan is an anionic heteropolysaccharide with a linear structure [88]. Fucoïdan from *Fucus spiralis* was obtained after the separation of polyphenols with a yield ($1.35 \pm 0.1\%$ dry seaweed) similar to values reported for brown macroalgae through conventional methods [46] and green methods [36]. The algal polymer extracted from brown species *Sargassum crispifolium* with 80% carbohydrate content by conventional isolation method and using distilled water as solvent generated a yield value of 1.50% dry basis [89]. Using a much higher extraction time (8 h), fucoïdan was separated from *Sargassum mcclurei*, *Sargassum polycystum* and *Turbinara ornata* using CaCl₂ 2%, with yields between 2.10 and 2.75% dried seaweed [90]. However, a separation method that increases the extraction yield is not necessarily favourable for structural integrity, purity and bioactive properties of the extracts [46].

In terms of chemical composition, elemental analysis of the E₂ extract (Table 2) indicates the presence of sulphur, which pinpoints the fucoïdan compound, as well as a high level of carbon (32.1%). Elemental assays in *Fucus serratus* and *Fucus vesiculosus* identified lower carbon values (21–28%) [91]. The amount of sulphur is much lower than that obtained from *Fucus vesiculosus* (9–12%) [34]. The H% content of the obtained fucoïdan is similar to the values reported for other brown macroalgae: 4.77% for *Fucus vesiculosus*, 4.78% for *Fucus serratus* [92], 4.64% for *Laminaria japonica* [93] and 4.99% for *Undaria pinnatifida* [94]. Moreover, a 3.47% hydrogen content reported for fucoïdan standard [95].

Table 2. Elemental analysis (CHNS) of fucoïdan and alginate extracts isolated from *Fucus spiralis* biomass.

Element	Fucoïdan	Alginate
C %	32.1 ± 0.4	31.4 ± 0.6
H %	4.0 ± 0.1	5.2 ± 0.3
S %	0.4 ± 0.0	0.3 ± 0.0
N %	ND	3.2 ± 0.2

E₂ was also subjected to FTIR characterisation (Figure 2). The spectra indicate the presence of carboxyl and carbonyl groups according to the peak registered at 1718 cm^{-1} [96]. Furthermore, an intense peak highlights the presence of the sulphate functional groups (S=O) characteristic of fucoïdan at a wavelength of 1226 cm^{-1} [34,97]. Two additional groups, one characteristic of sulphonamides, i.e., N=S=O [98], and one associated with C-O-SO₃, were observed at 1136 cm^{-1} and at 1024 cm^{-1} , respectively. Groups containing sulphur (S) play an important role in determining the bioactive properties of fucoïdan [67]. The sulphate functional group of marine macroalgae polysaccharides is associated with the antiviral properties of these compounds [88]. The peak at 1535 cm^{-1} indicates the presence of an N-H protein group in the structure of the amide II. The peak identified at 1408 cm^{-1} was assigned to the C-N group of amide III, whereas that identified at 1613 cm^{-1} was assigned to the C=O functional group of amide I or to the functional groups

of uronic acids [98]. Additionally, wavelengths 812 and 886 cm^{-1} were associated with units of mannuronic acid in the structure of alginate [97,99], which indicates that a complete separation of fucoidan could not be achieved.

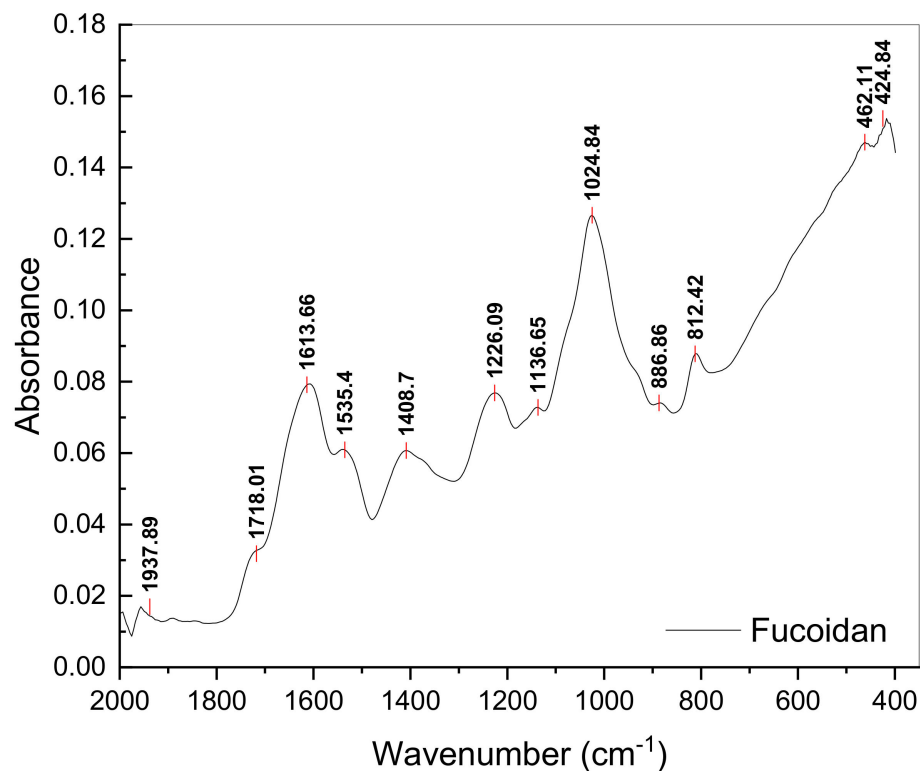


Figure 2. FTIR spectrum of fucoidan extract separated from *Fucus spiralis* biomass.

The NMR spectrum (Figure 3) of E_2 highlights the chemical structures of fucoidan. The peak at 5.337 ppm was assigned to α 3-linked 2-mono-*O*-sulfated-L-fucopyranose residues [100]. Another characteristic structure, - α -L-fucose, was identified through the intense signal at 4.8 ppm. The several intense peaks at 3.68–3.88 ppm were attributed to 4-linked 2-mono-*O*-sulfated L-fucopyranose residues [100], whereas the peak at 3.37 ppm was associated with β -d-xylose [98]. Finally, the signal at 1.26 was assigned to the C6 methyl proton group of L-fucopyranose [100,101].

Only traces of mono- or oligosaccharides in the supernatant and in the residual wash waters obtained from the separation of fucoidan extract were determined. Considering that the acid extraction of fucoidan may also remove any remaining polyphenols [102], the TPC of the supernatant was analysed. The obtained result, 1.68 mg GAE g^{-1} , indicates that following fucoidan extraction, a high polyphenolic content is still released equal to more than half of the total polyphenolic content initially separated initially UAE. Some studies have associated the antioxidant activity of fucoidan extracts with the presence of polyphenolic impurities [37].

3.3.3. Yield and Chemical Characterization of Alginate Extract (E_3)

Alginate is the third targeted bioproduct and was separated with a yield of $47.0 \pm 1.0\%$ dry seaweed. This value is significantly higher than the alginate yield obtained from *Fucus vesiculosus* ($16.2 \pm 3.2\%$ *w/w*) [103]. In a study on *Durvillaea potatorum* an alginate yield of 36.55% *w/w* was reported [104]. Other authors reported a 21.06% *w/w* yield for alginate separated from *Laminaria japonica* using Na_2CO_3 2% for 5 h at 60 °C [41], and a similar value was reported for *Ecklonia radiata* using Na_2CO_3 0.2 M for 120 min at 45 °C [46].

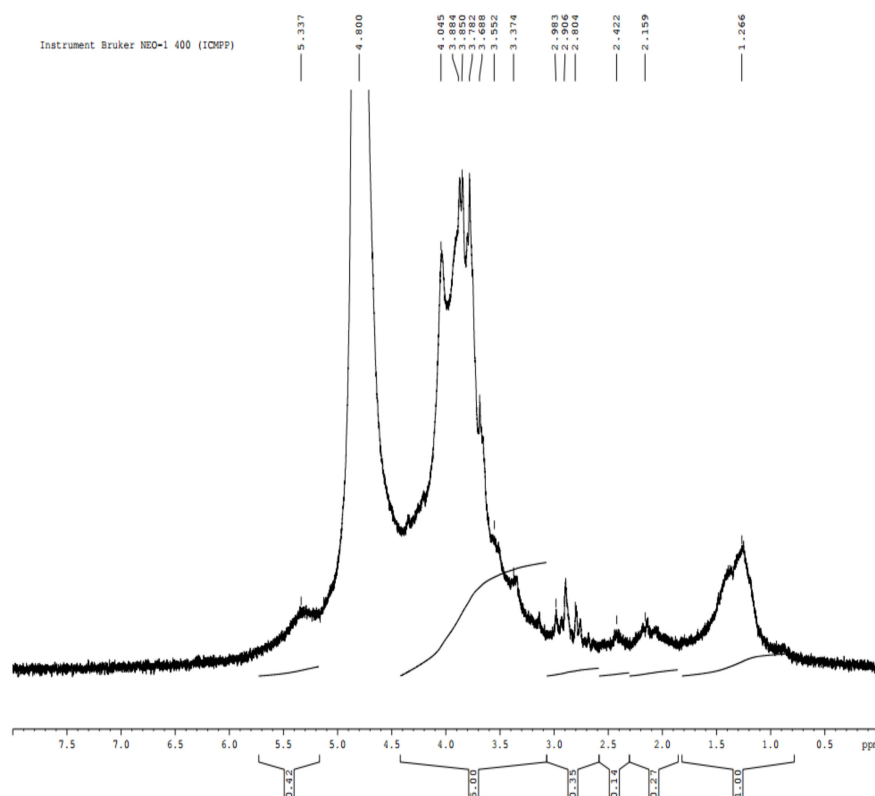


Figure 3. NMR spectrum of fucoidan extract separated from *Fucus spiralis* biomass.

In terms of chemical composition, the elemental analysis of E₃ (Table 2) shows similar carbon and hydrogen percentage values to those reported in the literature [105]. The amount of sulphur is lower than the alginate fraction from *Sargassum fusiforme* (12.5%) [106]. The presence of sulphur in the composition of the alginate extract can be beneficial for bioactive properties, but it also indicates fucoidan residues.

Alginate is a marine polysaccharide found in macroalgae, as well as other marine organisms. Its structure comprises homogenous or heterogenous disposition of β-D-mannuronic acid (M) and α-L-guluronic acid (G) [107]. Spectral analysis of E₃ (Figure 4) identified the presence of guluronic units (G) at 1017 cm⁻¹ [96,97,99] and mannuronic units (M) at 872 and 812 cm⁻¹ [97,99], confirming the isolation of alginate. The presence of uronic acids is also indicated by the functional groups identified at 1598 and 1397 cm⁻¹. Furthermore, sulphate (characteristic of fucoidan) was observed at 1159 and 1248 cm⁻¹ [97], suggesting that the extraction was not completely selective.

The ¹H NMR spectrum of E₃ representing the uronic acid sequence is depicted in Figure 5. The peak at 5.03 ppm was associated with H¹ of α-L-guluronic acid [106,108]. The presence of M units in the extract was indicated through an intense peak at 4.62 ppm, which was assigned to β-d-mannuronic acid residues [106,108]. Resonance signals of G units were also identified by peaks at 4.45 and 4.3 ppm [109].

In the supernatant resulting after E₃ separation, a total polyphenolic content (TPC) of 0.18 mg GAE g⁻¹ was determined. The presence of polyphenols is due to the use of ethanol, which facilitates the release of residual aromatic fractions from the cell wall of the macroalgae [19].

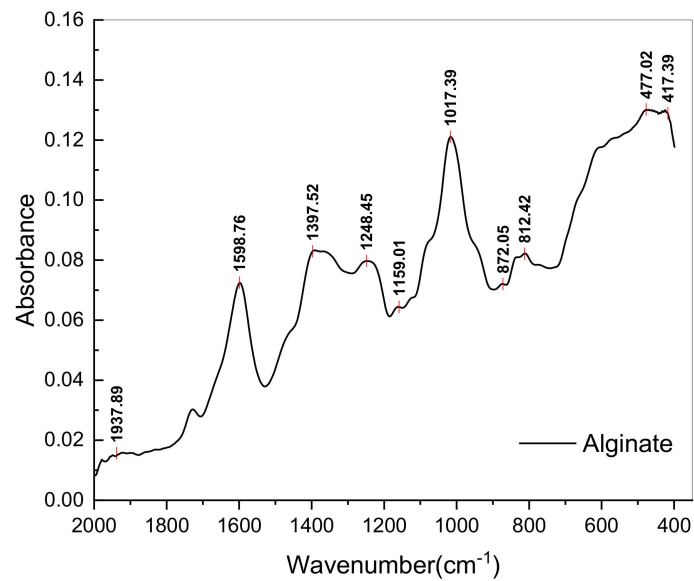


Figure 4. FTIR spectrum of alginate extract separated from *Fucus spiralis* biomass.

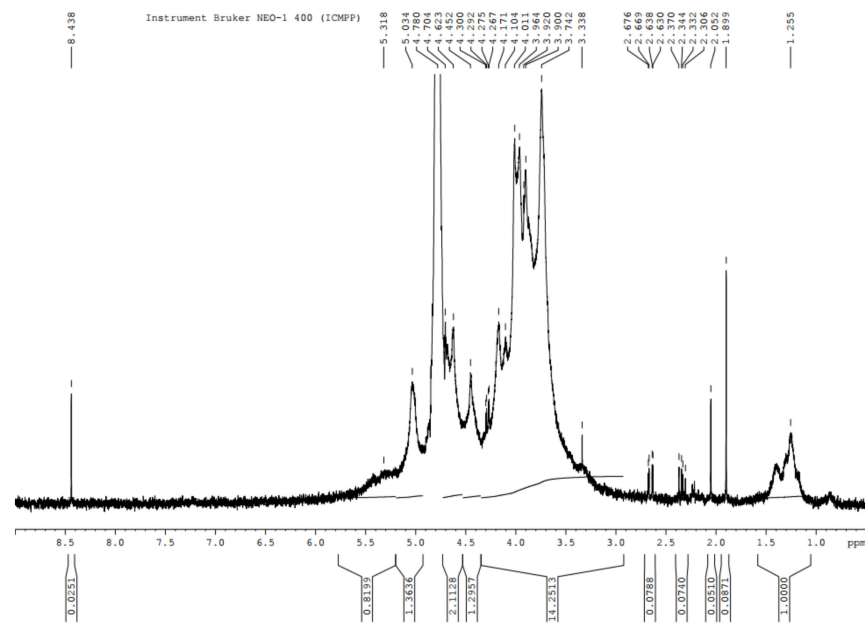


Figure 5. NMR spectrum of alginate extract separated from *Fucus spiralis* biomass.

3.4. Assessment of Targeted Molecule Bioactivity

3.4.1. Electrochemical DNA Binding Study

In the absence of DNA, the voltammograms of compounds E₂ and E₃ in a phosphate-buffered solution (KH₂PO₄/K₂HPO₄) show a cathodic peak potential at −94 and −342 mV, respectively. On the other hand, in the presence of a concentration of 6.88 μM of DNA, the peak potential of E₂ was shifted to a more negative value of −173 mV, with an apparently negative shift of −79 mV, whereas the peak potential of E₃ was shifted to a more positive value of −335 mV, with an apparently positive shift of +7 mV (Figure 6). These results indicate that E₂ interacts with DNA via an electrostatic mode [55,56], whereas the obvious positive shifts of peak potentials of E₃ indicate that the interaction mode may be intercalation between E₃ and DNA. The voltammograms also show a drop in the cathodic peak current densities, which can be attributed to slow diffusion of the formed DNA-E₂ and DNA-E₃ adducts.

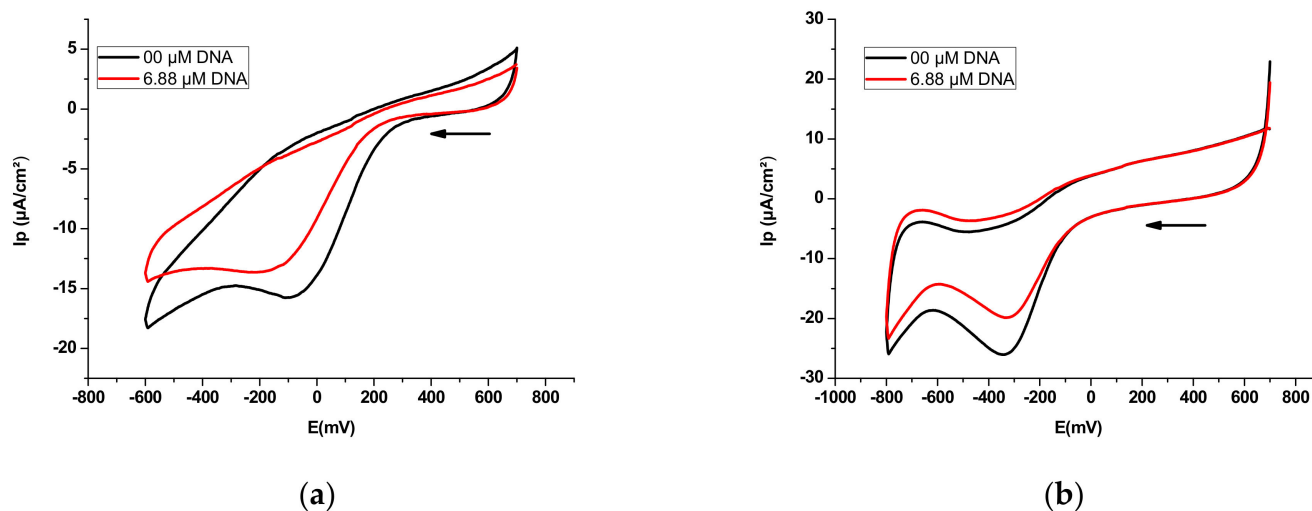


Figure 6. Cyclic voltammograms of polysaccharidic extract in 12 mL of 0.1 M phosphate-buffered solution ($\text{KH}_2\text{PO}_4/\text{K}_2\text{HPO}_4$, pH = 7.2) recorded at 0.1 V s^{-1} potential sweep rate on a platinum disk electrode at 298 K in the absence and presence of $6.88 \mu\text{M}$ DNA with supporting electrolyte 0.1 M Bu_4NBF_4 : (a) extract E_2 (65 mg); (b) extract E_3 (60 mg).

The negative shift in the cathodic peak potential that occurred as a result of the addition of DNA to E_2 was caused by the electrostatic interaction of the anionic drug with the DNA backbone [56,110]; therefore, the obvious negative peak potential shift (cathodic shift) in the CV behaviour of E_2 with the addition of DNA can be attributed to the electrostatic interaction (H-bonding) between E_2 and DNA. This negative peak potential shift further indicates that E_2 residue is easier to reduce in the presence of a negative DNA environment. However, the positive shift observed in the cathodic peak potential caused by the addition of DNA to E_3 can be explained by the intercalation interaction mode between E_3 and DNA. This positive shift further indicates that the formed adduct DNA- E_3 is electrochemically more stable than E_3 .

The binding constant of the interactions of E_2 and E_3 with DNA can be calculated based on the decrease in the cathodic peak current density of DNA- E_2 and DNA- E_3 adducts relative to free E_2 and E_3 , respectively, using the following Equation (2) [111]:

$$\log \frac{1}{[\text{DNA}]} = \log K_b + \log \frac{i_p}{i_{p_0} - i_p} \quad (2)$$

where $[\text{DNA}]$ is the DNA concentration (M); K_b represents the binding constant (M^{-1}); and i_{p_0} and i_p denote the cathodic peak current density of the free and DNA-bound compound ($\mu\text{A}\cdot\text{cm}^{-2}$), respectively.

The plot of $\log 1/[\text{DNA}]$ versus $\log i_p/i_{p_0} - i_p$ should afford a straight line with a 'y' intercept equal to the logarithm of binding constant K_b . To obtain the plot $\log 1/[\text{DNA}]$ versus $\log i_p/i_{p_0} - i_p$, voltammograms of E_2 and E_3 were obtained in the absence and presence of an increasing concentration of DNA (Figure 7).

The plot of $\log 1/[\text{DNA}]$ versus $\log i_p/i_{p_0} - i_p$ is shown in Figure 8, which is a straight line with a 'y' intercept equal to the logarithm of binding constant K_b .

The values of binding constants obtained from the regression of Figure 8 are tabulated in Table 3.

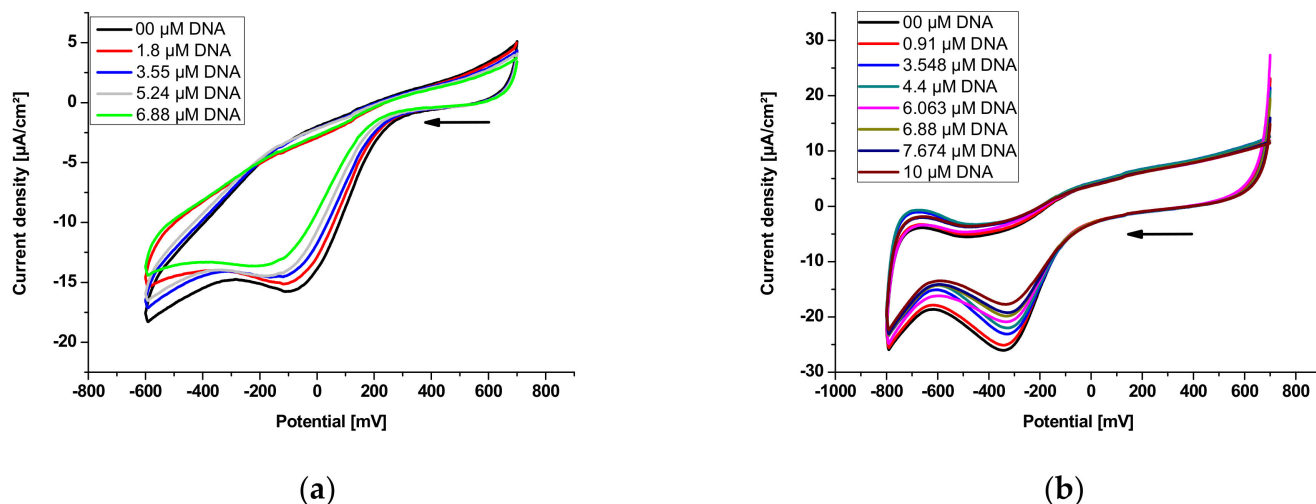


Figure 7. Cyclic voltammograms of polysaccharidic extracts in 12 mL of 0.1 M phosphate-buffered solution ($\text{KH}_2\text{PO}_4/\text{K}_2\text{HPO}_4$, pH = 7.2) recorded at a 0.1 Vs^{-1} potential sweep rate on a platinum disk electrode at 298 K in the absence and presence of an increasing concentration of DNA with supporting electrolyte 0.1 M Bu_4NBF_4 : (a) E_2 (65 mg); (b) E_3 (60 mg).

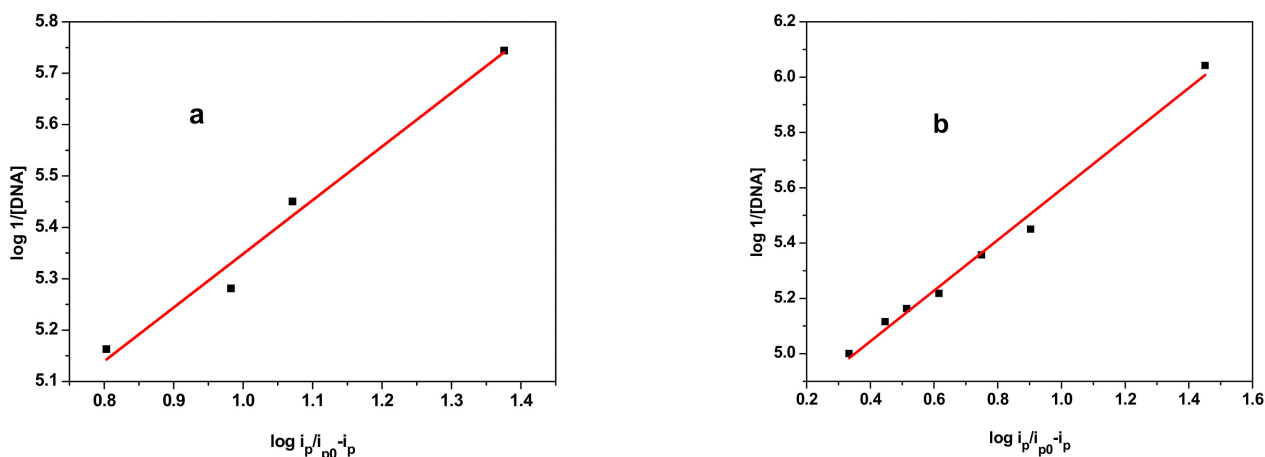


Figure 8. Plots of $\log(1/(1 - (i_p/i_{p0})))$ versus $\log 1/[\text{DNA}]$ used to calculate the binding constants of the E_2 and E_3 samples with DNA: (a) E_2 ; (b) E_3 .

The binding free energy change was calculated using the following Equation (3) [112]:

$$\Delta G = -RT \ln K_b \tag{3}$$

where ΔG is the binding free energy in $\text{KJ}\cdot\text{mol}^{-1}$, R is the gas constant ($8.32 \text{ J}\cdot\text{mol}^{-1}\cdot\text{K}^{-1}$) and T is the absolute temperature (298 K). Values are also listed in Table 3.

Table 3. Binding constants and binding free-energy values for the interactions of E_2 and E_3 with DNA based on CV data presented in Figure 8.

Sample	Equation	R^2	$K (\text{M}^{-1})$	$-\Delta G (\text{KJ}\cdot\text{mol}^{-1})$
E_2	$y = 1.0445x + 4.3039$	0.972	2.01×10^4	24.57
E_3	$y = 0.9162x + 4.6782$	0.989	4.77×10^4	26.71

The order of the magnitude of binding site size (s) in terms of the base pair of DNA can provide information about the interaction of small molecules with DNA. The binding site size(s) can be calculated according to the following Equation (4) [113]:

$$\frac{C_b}{C_f} = K_b \left(\frac{\text{free base pairs}}{s} \right) \quad (4)$$

where s represents the binding site size in terms of the base pair, K_b is the binding constant, C_f is the concentration of free compound and C_b represents the concentration of DNA-bound compound.

Considering the concentration of DNA in terms of nucleotide phosphate (NP), the concentration of the DNA base pair is expressed as $[DNA]/2$, so Equation (5) can be expressed as:

$$\frac{C_b}{C_f} = K_b \left(\frac{[DNA]}{2s} \right) \quad (5)$$

The C_b/C_f ratio is equal to $(i_{p0} - i_p)/i_p$ [114], which are the values of the experimental peak current densities.

The obtained values of binding site size for both samples are summarized in Table 4. The low value of the binding site size of E_2 further indicates the electrostatic interaction of E_2 with DNA. In addition, the relatively higher value of the binding site size of E_3 confirms the intercalation interaction of E_3 with DNA.

Table 4. Values of binding site size obtained using the plot of C_b/C_f versus (DNA).

Adduct	Equation	R ²	s
DNA- E_2	$y = 0.02193x - 8.287 \times 10^{-5}$	0.828	0.02
DNA- E_3	$y = 0.0486x - 0.027$	0.987	0.12

The diffusion coefficient is a parameter that further confirms the interaction of E_2 or E_3 with DNA; free E_2 or E_3 diffuses more rapidly in solution than their adducts, E_2 -DNA or E_3 -DNA. The diffusion coefficients of the free and DNA-bound forms of E_2 and E_3 were determined using the Randles–Sevcik Equation (6) [115].

$$i = 2.69 \times 10^5 n^{\frac{3}{2}} A C D^{\frac{1}{2}} v^{\frac{1}{2}} \quad (6)$$

where n is the number of electrons per species reaction, C is the bulk concentration ($\text{mol}\cdot\text{cm}^{-3}$) of the electroactive species, D is the diffusion coefficient (cm^2/s) and v is the scan rate (V/s).

The plots of \sqrt{v} versus i_p are displayed in Figure 9, whereas the diffusion coefficient values are presented in Table 5. The diffusion coefficients of the DNA- E_2 and DNA- E_3 adducts are lower than those for the free E_2 and E_3 samples, indicating the formation of a high-molecular-weight complex that diffuses slowly toward the electrode.

Table 5. Diffusion coefficient values of the free and DNA-bound forms of E_2 and E_3 .

Compound	Equation	R ²	D (cm^2/s)
E_2	$y = -1.29665x - 4.16404$	0.938	1.161×10^{-7}
DNA- E_2	$y = -1.31838x + 0.02649$	0.968	0.721×10^{-7}
E_3	$y = -2.9437x + 1.3783$	0.972	7.086×10^{-9}
DNA- E_3	$y = -2.6593x + 8.1874$	0.992	5.783×10^{-9}

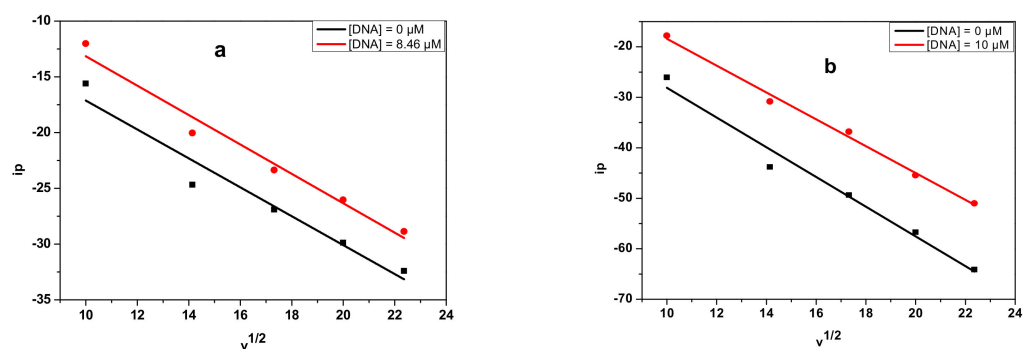


Figure 9. Plots of \sqrt{v} versus i_p used to calculate the coefficient diffusion of the free and E_2 - and E_3 -bound DNA: (a) E_2 ; (b) E_3 .

3.4.2. Electronic Spectroscopy DNA Binding Study

The absorption spectra of E_2 and E_3 in the absence and presence of a gradually increasing concentration of DNA stock solution are shown in Figure 10. In the ultraviolet region, E_2 has one absorption peak at 281 nm (Figure 10a), and E_3 also displays one peak at 268 nm (Figure 10b).

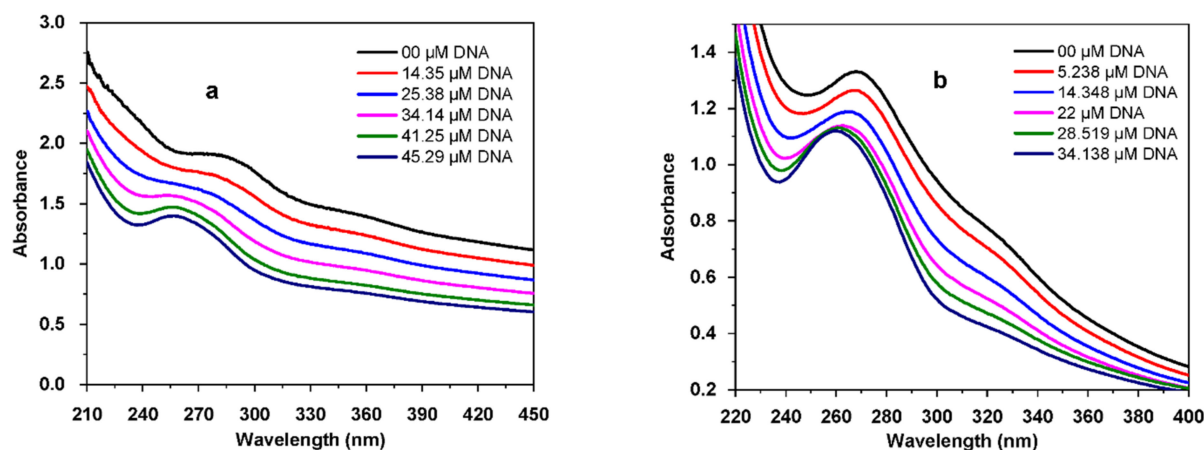


Figure 10. UV-visible absorption spectra of polysaccharidic extracts in the presence of increasing concentrations of DNA in 0.1 M phosphate-buffered solution ($\text{KH}_2\text{PO}_4/\text{K}_2\text{HPO}_4$) at pH = 7.2 and 298 K: (a) 2 mg of E_2 ; (b) 6 mg of E_3 .

Upon addition of DNA, a significant hypochromicity was observed without any noticeable shift in the position of the maximum absorption peak, clearly indicating the formation of the adducts E_2 -DNA and E_3 -DNA, with hypochromicity further suggesting the groove-binding property of E_2 and E_3 compounds with respect to the double-stranded DNA [116].

Absorbance values were changed by increasing DNA concentration to evaluate the binding constant using the following Equation (7), [117]:

$$\frac{A_0}{A - A_0} = \frac{\varepsilon_G}{\varepsilon_{H-G} - \varepsilon_G} + \frac{\varepsilon_G}{\varepsilon_{H-G} - \varepsilon_G} \frac{1}{K_b [DNA]} \quad (7)$$

where $[DNA]$ is the DNA concentration; K_b is the binding constant; A_0 and A are the absorbance of E_2 and E_3 , respectively, in the absence and presence of DNA; and ε_G and ε_{H-G} are their extinction coefficients, respectively.

The binding constant (K_b) is obtained from the intercept to slope ratio of the plot of $A_0/(A - A_0)$ versus $1/[DNA]$ (Figure 11). The binding free-energy change was calculated using Equation (4). Values for both binding constant and free energy are listed in Table 6.

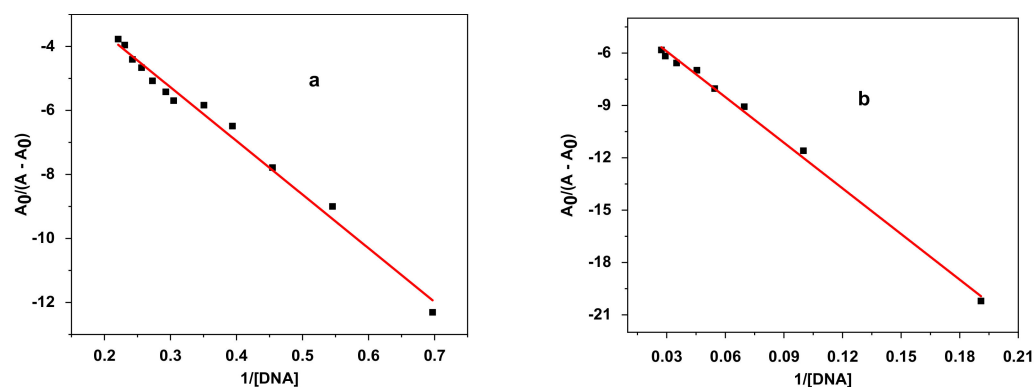


Figure 11. Plots of $A_0/(A - A_0)$ versus $1/[DNA]$ used to calculate the binding constants of algal polysaccharidic extracts with DNA: (a) E_2 and (b) E_3 .

Table 6. Binding constants and binding free energy values for the interaction of algae polysaccharidic extracts with DNA from the UV data shown in Figure 6: E_2 and E_3 .

Sample	Equation	R^2	$K (M^{-1})$	$-\Delta G (KJ \cdot mol^{-1})$
E_2	$y = -16.76582x - 0.24616$	0.986	1.47×10^4	23.79
E_3	$y = -87.1888x - 3.2914$	0.998	3.78×10^4	26.13

The values of the binding free energy and binding constant obtained from cyclic voltammetry (Table 3) and electronic spectroscopy (Table 6) are in agreement.

3.4.3. Superoxide Anion Radical Assay

The one-electron reduction in molecular oxygen generates the superoxide anion radical; this reactive oxygen species causes cell degradation and leads to aging and various diseases [118]. Antioxidants of various natures play a major role in the prevention of these occurrences. Figure 12 shows the decrease in the anodic peak potential of the voltammograms of the radical $O_2^{\cdot -}$ in the presences of varying concentrations of E_2 and E_3 .

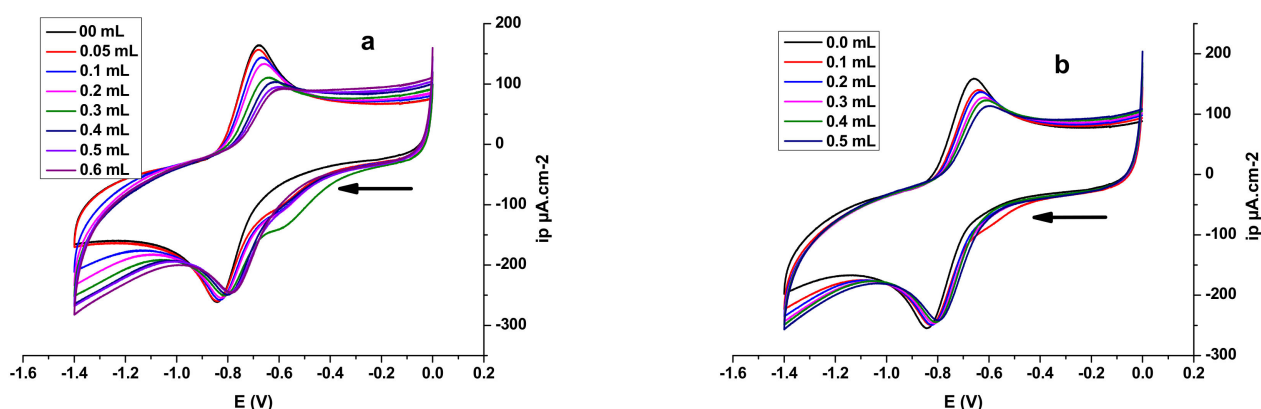


Figure 12. Cyclic voltammograms of oxygen-saturated DMSO/0.1 M Bu_4NBF_4 on a Pt electrode in the absence and presence of varying concentrations of algal polysaccharidic extracts at a scan rate 100 mV/s: (a) E_2 ; (b) E_3 .

The equations obtained from the linear calibration graph in the studied concentration range for E_2 , E_3 and α -tocopherol are summarized in Table 7 (where y represents the value of the anodic peak current density of $O_2^{\cdot -}$, and x represents the value of the sample concentrations, expressed as mg/mL). All the tests were performed in triplicate, and the graph was plotted according to the averages of three observations. The obtained results

indicate that the $O_2^{\cdot-}$ radical-scavenging activity of E_2 is slightly higher than that E_3 , and both are higher than that of the standard antioxidant α -tocopherol.

Table 7. IC_{50} values (mg/mL) obtained using $O_2^{\cdot-}$ radical-scavenging activity.

Sample	Equation	R^2	IC_{50}
E_2	$y = 3.21057x + 0.08397$	0.918	0.13 ± 0.04
E_3	$y = 2.3336x + 0.0658$	0.984	0.19 ± 0.06
α -tocopherol	$y = 15.99x + 1.37$	0.950	3.04 ± 0.30

In the presence of E_2 and E_3 , the anodic peak potentials of the redox couple $O_2/O_2^{\cdot-}$ were shifted to more negative potentials. This shift was associated with a significant decrease in anodic peak current density. The significant drop in anodic peak current density can be assigned to the decrease in $O_2^{\cdot-}$ concentration due to the formation of slowly diffusing $E_2-O_2^{\cdot-}$ and $E_3-O_2^{\cdot-}$ complexes. Typical CV behaviour of $O_2^{\cdot-}$ in DMSO/0.1 M Bu_4NBF_4 in the potential window of 0.0 to -1.4 V at a Pt electrode in the presence of E_2 (a) and E_3 (b) is shown in Figure 13.

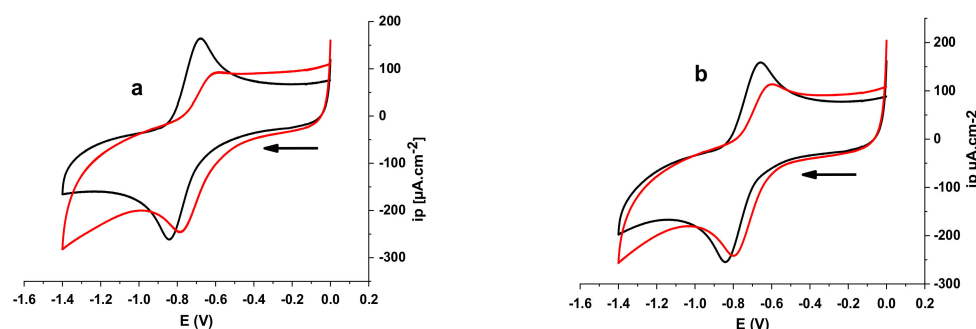


Figure 13. Cyclic voltammograms of $O_2^{\cdot-}$ on a polished Pt electrode in the absence (black line) and presence of algal polysaccharidic extracts in DMSO solution with supporting electrolyte 0.1 M Bu_4NBF_4 at $100 \text{ mV}\cdot\text{s}^{-1}$: (a) 0.6 mL of E_2 ; (b) 0.5 mL of E_3 .

The free $O_2/O_2^{\cdot-}$ redox couple exhibits one oxidation peak at -0.6764 V and one reduction peak at -0.8411 V. Figure 14 also shows the effect of the addition of E_2 and E_3 to a solution of $O_2^{\cdot-}$ in DMSO/0.1M Bu_4NBF_4 on the oxidation peak current density of the $O_2/O_2^{\cdot-}$ couple. The decrease in the anodic peak current density caused by the addition of E_2 or E_3 can be explained by the reaction of $O_2^{\cdot-}$ with E_2 and E_3 [119]. This decrease can be used to calculate the binding constant, whereas the shift in peak potential values can be exploited to determine the mode of interaction [120,121].

The addition of varying concentrations of E_2 or E_3 in DMSO to a solution of DMSO saturated with commercial oxygen provokes a remarkable decrease in the peak current density (Figure 13). The substantial diminution in peak current density can be attributed to the decrease in free $O_2/O_2^{\cdot-}$ radical concentration due to the formation of $E_2-O_2^{\cdot-}$ or $E_3-O_2^{\cdot-}$ product. The gradual decrease in peak current density of the $O_2/O_2^{\cdot-}$ redox couple by increasing E_2 or E_3 concentrations can be exploited to calculate the binding constant by applying the following Equation (8) [122]:

$$\log\left(\frac{1}{C}\right) = \log(K_b) + \log\left(\frac{i_{p_a}}{i_{p_{a0}} - i_{p_a}}\right) \quad (8)$$

where $i_{p_{a0}}$ and i_{p_a} are the peak currents of the superoxide anion radical in the absence and presence of E_2 and E_3 , respectively; whereas C is the concentration of E_2 or E_3 . As C is not known, this term was replaced by the volume of the added E_2 or E_3 (ΔV).

The volume of the solution containing $O_2^{\cdot-}$ is fixed; thus, the addition of volume increments of E_2 or E_3 is proportional to the addition of an increased number of moles (i.e., concentration) of the compounds. Binding free energy in this case can be calculated using the following Equation (9) [119]:

$$\log\left(\frac{1}{\Delta V}\right) = \log(K_b) + \log\left(\frac{i_{pa}}{i_{pa0} - i_{pa}}\right) \quad (9)$$

In Equation (9), ΔV represents the added volume of E_2 or E_3 (mL); K_b refers to the binding constant; and i_{pa0} and i_{pa} are the anodic peak current densities in the absence and presence of E_2 or E_3 , respectively.

The binding free-energy change was calculated using Equation (4). The values of binding constant and binding free energy are summarized in Table 8.

Table 8. Binding constant values for E_2 and E_3 with $O_2^{\cdot-}$ from CV data at $T = 298$ K.

Compound	Equation	R^2	K (L^{-1})	$-\Delta G$ ($KJ \cdot mol^{-1}$)
$E_2-O_2^{\cdot-}$	$y = 0.96108x + 3.02683$	0.978	1.06×10^3	17.28
$E_3-O_2^{\cdot-}$	$y = 1.28666x + 2.55632$	0.891	0.36×10^3	14.59

The addition of 0.6 mL of E_2 or 0.5 mL of E_3 has caused a slight shift in peak potential ΔE^0 in the negative direction in association with a remarkable decrease in anodic peak current density due to the scavenging activity of the added compounds [123] (Figure 13). The significant drop in anodic peak current density can be assigned to the decrease in $O_2^{\cdot-}$ radical concentration due to the formation of $E_2-O_2^{\cdot-}$ or $E_3-O_2^{\cdot-}$ complexes. The peak potential shift of the $O_2/O_2^{\cdot-}$ redox couple in the negative direction in the presence of E_2 or E_3 indicates that the oxidation of $O_2^{\cdot-}$ occurs more easily in the presence of E_2 or E_3 because its oxidized form, O_2 , is less strongly attached to E_2 or E_3 than its reduced form, $O_2^{\cdot-}$. For such a system, where both forms of the $O_2/O_2^{\cdot-}$ redox couple interact with E_2 or E_3 , Figure 14 can be applied [114].

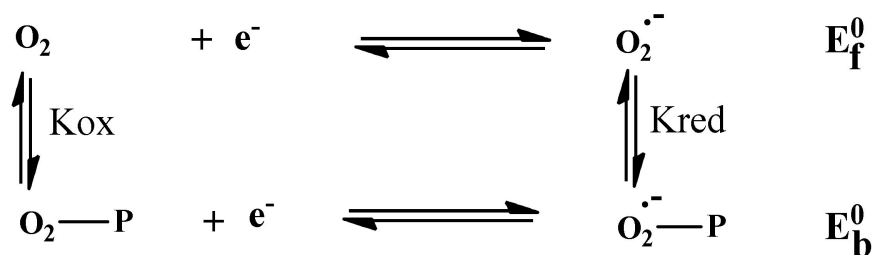


Figure 14. Redox process of the free and E-bound $O_2/O_2^{\cdot-}$ redox couple, where E represents E_2 or E_3 .

The application of the Nernst relation to the process presented in Figure 14 leads to Equation (10) [124]:

$$\Delta E^0 = E_b^0 - E_f^0 = E^0(O_2^{\cdot-} - P) - E^0(O_2^{\cdot-}) = 0.059 \log \frac{K_{red}}{K_{ox}} \quad (10)$$

where E_f^0 and E_b^0 are the formal potentials of the $O_2/O_2^{\cdot-}$ couple in the free and bound forms, respectively.

The ratio of the binding constants is calculated by replacing ΔE^0 from Table 9 in Equation (10).

Table 9. Electrochemical data of free and $O_2^{\cdot-}$ -bound forms of E_2 and E_3 used to calculate ratio of binding constants.

Compound	E_{pa} (V)	E_{pc} (V)	E_0 (V)	$E_b^0 - E_f^0$ (V)	K_{red}/K_{ox}
E_2	-0.6764	-0.8411	-0.75875	0.06765	14.01
$E_2-O_2^{\cdot-}$	-0.6018	-0.7804	-0.6911		
E_3	-0.6575	-0.8411	-0.7493	0.05095	7.03
$E_3-O_2^{\cdot-}$	-0.6018	-0.7949	-0.69835		

The obtained ratios of the binding constants indicate that the reaction of E_2 with $O_2^{\cdot-}$ is twice as strong as that of E_3 with $O_2^{\cdot-}$, which explains the high antioxidant activity of E_2 . Furthermore, values of the ratio of the binding constants indicate that the interaction of the reduced form ($O_2^{\cdot-}$) of the couple $O_2/O_2^{\cdot-}$ with E_2 or E_3 is always stronger than that of the oxidized form (O_2).

3.4.4. DPPH (2,2-Diphenyl-1-picrylhydrazyl) Assay

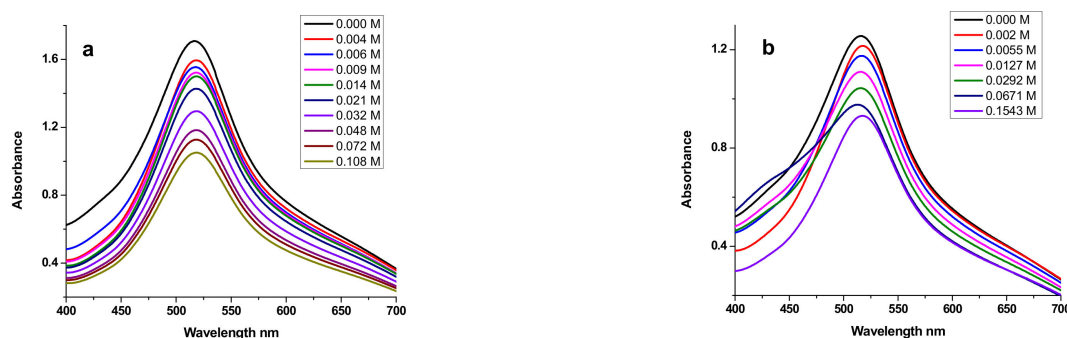
The antioxidant activity of the macroalgae extracts was also evaluated using DPPH. The equations obtained from the linear calibration graph in the studied concentration range for E_2 and E_3 samples and α -tocopherol are summarized in Table 10 (where y represents the value of the absorbance of DPPH, and x represents the value of sample concentrations, expressed as mg/mL). All tests were performed in triplicate, and the graph was plotted with the averages of three observations.

Table 10. IC_{50} values (mg/mL) obtained using DPPH radical-scavenging activity.

Sample	Equation	R^2	IC_{50}
E_2	$y = 0.114x - 0.0123$	0.997	4.49 ± 0.21
E_3	$y = 0.1051x - 0.0081$	0.998	4.83 ± 0.34
α -tocopherol	$y = 1.5646x + 29.093$	0.996	4.81 ± 0.32

The obtained results presented in Table 10 indicate that the DPPH radical-scavenging activity of E_2 is slightly higher than that of E_3 , with both approximately equivalent to that of the standard antioxidant α -tocopherol.

The independence of absorbance upon addition of the gradually increasing concentration of E_2 and E_3 to a DPPH solution is presented in Figure 15. The figure clearly shows a decrease in absorbance following the addition of the studied compounds.

**Figure 15.** UV-visible absorption spectra of a solution of DPPH (2 mg) in 10 mL acetonitrile in the presence of increasing concentrations of polysaccharidic extracts in acetonitrile at 298 K: (a) E_2 ; (b) E_3 .

The change in absorbance values by increasing E_2 and E_3 concentration was used to evaluate the intrinsic binding constant by employing Equation (11), [125].

$$\frac{A_0}{A - A_0} = \frac{\varepsilon_G}{\varepsilon_{H-G} - \varepsilon_G} + \frac{\varepsilon_G}{\varepsilon_{H-G} - \varepsilon_G} \frac{1}{K_b [P]} \quad (11)$$

where $[P]$ is the E_2 or E_3 concentration; K_b is the binding constant; A_0 and A are the absorbance values of DPPH in the absence and presence of E_2 or E_3 , respectively; and ε_G and ε_{H-G} are their extinction coefficients, respectively. The constant K_b is obtained from the intercept-to-slope ratio of the plot of $A_0/(A - A_0)$ versus $1/[P]$ (Figure 16).

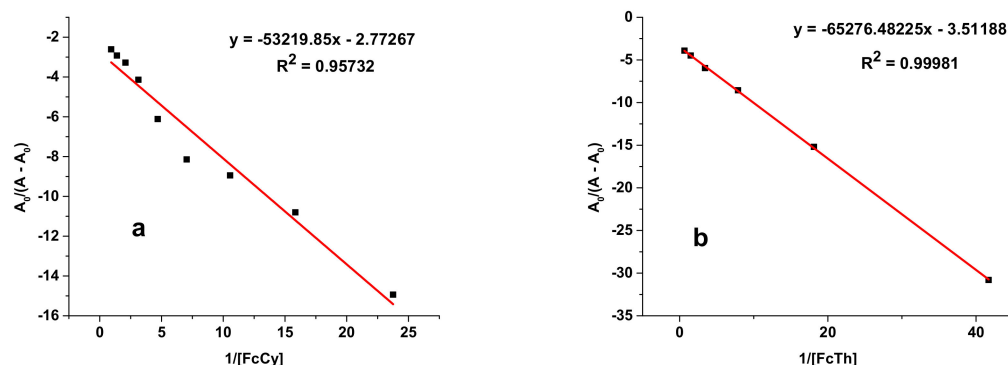


Figure 16. Plots of $A_0/(A - A_0)$ versus $1/[P]$ used to calculate the binding constants of algal compounds with DPPH: (a) E_2 ; (b) E_3 .

The binding free-energy change was calculated using Equation (4), and the values are presented in Table 11.

Table 11. Binding constant and binding free-energy values for E_2 and E_3 with DPPH from UV data at 298 K.

Adduct	Equation	R^2	K (M^{-1})	$-\Delta G$ ($KJ \cdot mol^{-1}$)
E_2 -DPPH	$y = -53,219.85x - 2.77267$	0.957	52.1	9.80
E_3 -DPPH	$y = -65,276.482x - 3.5119$	0.999	53.8	9.88

Values of binding free energy listed in the previous table are very close to each other, which may indicate that the studied compounds bind to DPPH in the same manner via their nitrogen atoms.

4. Conclusions

In this study, we highlighted a biorefinery approach applied for the first time, at lab scale, to *Fucus spiralis* biomass, with the aim of generating high-value compounds with bioactive properties.

Chemical characterization showed the presence of the targeted algal polysaccharides in the obtained extracts through the identification of S=O (1226 cm^{-1}), N=S=O (1136 cm^{-1}) and C-O-SO₃ (1024 cm^{-1}) groups characteristic to fucoidan, guluronic, (G) (1017 cm^{-1}) and mannuronic units (M) (872 and 812 cm^{-1}) specific to the alginate compound.

Cyclic voltammetry and electronic spectroscopy were performed in order to determine the DNA binding affinity and antioxidant activity of E_2 (fucoidan) and E_3 (alginate) extracts. A comparison of the binding constant and free-energy values of the generated adducts indicated consistency in the generated results with respect to the interaction of fucoidan and alginate with DNA.

Both polysaccharides demonstrated antioxidant activity, with fucoidan having a slightly higher effect than alginate in both DPPH and $O_2^{\cdot -}$ radical assays. In DPPH as-

says, the registered IC₅₀ values of E₂ and E₃ were 4.49 ± 0.21 and 4.83 ± 0.34 mg/mL, respectively. However, in O₂^{•−} assays, the obtained IC₅₀ values were 0.13 ± 0.04 mg/mL for fucoxanthin and 0.19 ± 0.06 mg/mL for alginate. Both assays revealed higher antioxidant activity than that of the control antioxidant, α-tocopherol.

Author Contributions: Conceptualization, I.V. and T.L.; methodology, T.L. and E.L.; software, C.F. and E.L.; validation, I.V., T.L. and V.I.P.; formal analysis, C.F. and E.L.; investigation, C.F. and E.L.; resources, I.V., V.I.P. and T.L.; data curation, C.F. and E.L.; writing—original draft preparation, C.F.; writing—review and editing, I.V., T.L. and V.I.P.; visualization, C.F. and E.L.; supervision, I.V.; project administration, I.V.; funding acquisition, I.V. All authors have read and agreed to the published version of the manuscript.

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